

RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number:	09	19	11,132
Source:	01	PE	
Date Processed by STIC:	2	25	02

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.
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FOR CRF SUBMISSION QUESTIONS, PLEASE CONTACT MARK SPENCER, 703-308-4212.

FOR SEQUENCE RULES INTERPRETATION, PLEASE CONTACT ROBERT WAX, 703-308-4216. PATENTIN 2.1 e-mail help: patin21help@uspto.gov or phone 703-306-4119 (R. Wax) PATENTIN 3.0 e-mail help: patin3help@uspto.gov or phone 703-306-4119 (R. Wax)

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 3.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

http://www.uspto.gov/web/offices/pac/checker

4.

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail. Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom. Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

- 1. EFS-Bio (http://www.uspto.gov/ebc/efs/downloads/documents.htm, EFS Submission User Manual ePAVE)
- 2. U.S. Postal Service: U.S. Patent and Trademark Office, Box Sequence, P.O. Box 2327, Arlington, VA 22202
- Hand Carry directly to:
 U.S. Patent and Trademark Office, Technology Center 1600, Reception Area, 7th Floor, Examiner Name, Sequence Information, Crystal Mall One, 1911 South Clark Street, Arlington, VA 22202
 - U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
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Revised 01/29/2002



OIPE

RAW SEQUENCE LISTING DATE: 02/25/2002 PATENT APPLICATION: US/09/911,132 TIME: 10:07:43

Input Set : A:\seq listing.txt
Output Set: N:\CRF3\02252002\I911132.raw

Does Not Comply
Corrected Diskette Needed

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Errors on pp. 1-5.
      3 <110> APPLICANT: Roche Diagnostics GmbH
      5 <120> TITLE OF INVENTION: Expression of alkaline phosphatase in yeast
      7 <130> FILE REFERENCE: 5387/00/
C--> 9 <140> CURRENT APPLICATION NUMBER: US/09/911,132
C--> 10 <141> CURRENT FILING DATE: 2001-07-23
     12 <160> NUMBER OF SEQ ID NOS: 38
     14 <170> SOFTWARE: PatentIn Ver. 2.1
     16 <210> SEQ ID NO: 1
     17 <211> LENGTH: 1476
     18 <212> TYPE: DNA
     19 <213> ORGANISM: Bovine
     21 <400> SEQUENCE: 1
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     25 atgaatggca aactgggacc tgagacaccc ctggccatgg accagttccc atacgtggct 240
     26 ctgtccaaga catacaacgt ggacagacag gtgccagaca gcgcaggcac tgccactgcc 300
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     29 gcagggaagg ccgtgggagt ggtgaccacc accagggtgc agcatgcctc cccagccggg 480
     30 geetaegege acaeggtgaa eegaaaetgg taeteagaeg eegaeetgee tgetgatgea 540
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     43 taccggcagc aggcggccgt gcccctggct agcgagaccc acgggggcga agacgtggcg 1320
     44 gtgttcgcgc gaggcccgca ggcgcacctg gtgcacggcg tgcaggagga gaccttcgtg 1380
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     46 cccgccaccg ccaccagcat ccccgactag ggtacc
     48 <210> SEQ ID NO: 2
     49 <211> LENGTH: 40
                                                                           involid response, see
error summary
ficial shut, item 11.
     50 <212> TYPE: DNA
     51 <213> ORGANISM: Artificial Sequence
     53 <220> FEATURE:
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     54 <223> OTHER INFORMATION: Description of Artificial Sequence
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Input Set : A:\seq listing.txt

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60 <211> LENGTH: 36
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61 <212> TYPE: DNA
62 <213> ORGANISM: Artificial Sequence
64 <220> FEATURE:
65 <223> OTHER INFORMATION: Description of Artificial Sequence (Artificial
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72 <212> TYPE: PRT
73 <213> ORGANISM: Bovine
75 <400> SEQUENCE: 4
76 Leu Ile Pro Ala Glu Glu Glu Asn Pro Ala Phe Trp Asn Arg Gln Ala
                                        10
79 Ala Gln Ala Leu Asp Val Ala Lys Lys Leu Gln Pro Ile Gln Thr Ala
                                    25
                20
82 Ala Lys Asn Val Ile Leu Phe Leu Gly Asp Gly Met Gly Val Pro Thr
           35
                                40
85 Val Thr Ala Thr Arg Ile Leu Lys Gly Gln Met Asn Gly Lys Leu Gly
                            55
88 Pro Glu Thr Pro Leu Ala Met Asp Gln Phe Pro Tyr Val Ala Leu Ser
91 Lys Thr Tyr Asn Val Asp Arg Gln Val Pro Asp Ser Ala Gly Thr Ala
94 Thr Ala Tyr Leu Cys Gly Val Lys Gly Asn Tyr Arg Thr Ile Gly Val
                                   105
97 Ser Ala Ala Ala Arg Tyr Asn Gln Cys Asn Thr Thr Arg Gly Asn Glu
                               120
100 Val Thr Ser Val Ile Asn Arg Ala Lys Lys Ala Gly Lys Ala Val Gly
       130
                            135
                                                140
103 Val Val Thr Thr Arg Val Gln His Ala Ser Pro Ala Gly Ala Tyr
                        150
                                            155
106 Ala His Thr Val Asn Arg Asn Trp Tyr Ser Asp Ala Asp Leu Pro Ala
                    165
                                        170
109 Asp Ala Gln Lys Asn Gly Cys Gln Asp Ile Ala Ala Gln Leu Val Tyr
110
                180
                                    185
112 Asn Met Asp Ile Asp Val Ile Leu Gly Gly Arg Met Tyr Met Phe
           195
                                200
115 Pro Glu Gly Thr Pro Asp Pro Glu Tyr Pro Asp Asp Ala Ser Val Asn
                            215
                                                220
118 Gly Val Arg Lys Asp Lys Gln Asn Leu Val Gln Glu Trp Gln Ala Lys
                        230
                                            235
121 His Gln Gly Ala Gln Tyr Val Trp Asn Arg Thr Ala Leu Leu Gln Ala
                    245
                                        250
124 Ala Asp Asp Ser Ser Val Thr His Leu Met Gly Leu Phe Glu Pro Ala
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Input Set : A:\seq listing.txt

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127 Asp Met Lys Tyr Asn Val Gln Gln Asp His Thr Lys Asp Pro Thr Leu
            275
                                280
130 Ala Glu Met Thr Glu Ala Ala Leu Gln Val Leu Ser Arg Asn Pro Arg
131
        290
                            295
                                                 300
133 Gly Phe Tyr Leu Phe Val Glu Gly Gly Arg Ile Asp His Gly His His
134 305
                        310
                                             315
136 Asp Gly Lys Ala Tyr Met Ala Leu Thr Glu Ala Ile Met Phe Asp Asn
                    325
                                        330
139 Ala Ile Ala Lys Ala Asn Glu Leu Thr Ser Glu Leu Asp Thr Leu Ile
                                    345
140
                340
142 Leu Val Thr Ala Asp His Ser His Val Phe Ser Phe Gly Gly Tyr Thr
143
145 Leu Arg Gly Thr Ser Ile Phe Gly Leu Ala Pro Gly Lys Ala Leu Asp
        370
                            375
148 Ser Lys Ser Tyr Thr Ser Ile Leu Tyr Gly Asn Gly Pro Gly Tyr Ala
                        390
                                             395
151 Leu Gly Gly Gly Ser Arg Pro Asp Val Asn Gly Ser Thr Ser Glu Glu
                    405
                                        410
154 Pro Ser Tyr Arg Gln Gln Ala Ala Val Pro Leu Ala Ser Glu Thr His
155
                420
                                    425
                                                         430
157 Gly Gly Glu Asp Val Ala Val Phe Ala Arg Gly Pro Gln Ala His Leu
            435
                                440
160 Val His Gly Val Gln Glu Glu Thr Phe Val Ala His Ile Met Ala Phe
                            455
161
        450
                                                 460
163 Ala Gly Cys Val Glu Pro Tyr Thr Asp Cys Asn Leu Pro Ala Pro Ala
                                                                 480
                        470
                                             475
166 Thr Ala Thr Ser Ile Pro Asp
                    485
167
170 <210> SEQ ID NO: 5
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171 <211> LENGTH: 1476
172 <212> TYPE: DNA
173 <213> ORGANISM: Artificial Sequence
175 <220> FEATURE:
176 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
178 <400> SEQUENCE: 5
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180 gctttggatg ttgctaagaa gttgcaacca attcaaactg ctgctaagaa tgttattttg 120
181 tttttgggtg atggtatggg tgttccaact gttactgcta ctagaatttt gaagggtcaa 180
182 atgaatggta agttgggtcc agaaactcca ttggctatgg atcaatttcc atacgttgct 240
183 ttgtctaaga cttacaatgt tgatagacaa gttccagatt ctgctggtac tgctactgct 300
184 tacttgtgtg gtgttaaggg taattacaga actattggtg tttctgctgc tgctagatac 360
185 aatcaatgta atactactag aggtaatgaa gttacttctg ttattaatag agctaagaag 420
186 gctggtaagg ctgttggtgt tgttactact actagagttc aacatgcttc tccagctggt 480
187 gcttacgctc atactgttaa tagaaattgg tactctgatg ctgatttgcc agctgatgct 540
188 caaaagaatg gttgtcaaga tattgctgct caattggttt acaatatgga tattgatgtt 600
189 attttgggtg gtggtagaat gtacatgttt ccagaaggta ctccagatcc agaataccca 660
190 gatgatgctt ctgttaatgg tgttagaaag gataagcaaa atttggttca agaatggcaa 720
191 gctaagcatc aaggtgctca atatgtttgg aatagaactg ctttgttgca agctgctgat 780
192 gattctagtg ttactcattt gatgggtttg tttgaaccag ctgatatgaa gtataatgtt 840
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Input Set : A:\seq listing.txt

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193 caacaagatc atactaagga tccaactttg gctgaaatga ctgaagctgc tttgcaagtt 900
194 ttgtctagaa atccaagagg tttttacttg tttgttgaag gtggtagaat tgatcatggt 960
195 catcatgatg gtaaggctta tatggctttg actgaagcta ttatgtttga taatgctatt 1020
196 gctaaggcta atgaattgac ttctgaattg gatactttga ttttggttac tgctgatcat 1080
197 agtcatgttt tttcttttgg tggttacact ttgagaggta cttctatttt tggtttggct 1140
198 ccaggtaagg ctttggatag taagtettae aettetattt tgtatggtaa tggteeaggt 1200
199 tatgctttgg gtggtggttc tagaccagat gttaatggta gtactagtga agaaccatct 1260
200 tacagacaac aagctgctgt tccattggct agtgaaactc atggtggtga agatgttgct 1320
201 gtttttgcta gaggtccaca agctcatttg gttcatggtg ttcaagaaga aacttttgtt 1380
202 gctcatatta tggcttttgc tggttgtgtt gaaccataca ctgattgtaa tttgccagct 1440
203 ccagctactg ctactagtat tccagattaa ggtacc
                                                                       1476
205 <210> SEQ ID NO: 6
206 <211> LENGTH: 78
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207 <212> TYPE: DNA
208 <213> ORGANISM: Artificial Sequence
210 <220> FEATURE:
211 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
213 <400> SEQUENCE: 6
214 gcgcqaattc ttgattccag ctgaagaaga aaatccagct ttttggaata gacaagctgc 60
215 tcaagctttg gatgttgc
217 <210> SEQ ID NO: 7
218 <211> LENGTH: 70
219 <212> TYPE: DNA
220 <213> ORGANISM: Artificial Sequence
                                                                                Some
222 <220> FEATURE:
223 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
225 <400> SEQUENCE: 7
226 ccaaaaacaa aataacattc ttagcagcag tttgaattgg ttgcaacttc ttagcaacat 60
227 ccaaaqcttg
                                                                       70
229 <210> SEQ ID NO: 8
230 <211> LENGTH: 69
231 <212> TYPE: DNA
232 <213> ORGANISM: Artificial Sequence
                                                                               Serre
234 <220> FEATURE:
235 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
237 <400> SEQUENCE: 8
238 gaatgttatt ttgtttttgg gtgatggtat gggtgttcca actgttactg ctactagaat 60
239 tttgaaggg
241 <210> SEQ ID NO: 9
242 <211> LENGTH: 70
243 <212> TYPE: DNA
244 <213> ORGANISM: Artificial Sequence
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247 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
249 <400> SEQUENCE: 9
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253 <210> SEQ ID NO: 10
254 <211> LENGTH: 71
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Input Set : A:\seq listing.txt

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256 <213> ORGANISM: Artificial Sequence
                                                                              Barre
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259 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
261 <400> SEQUENCE: 10
262 gctatggatc aatttccata cgttgctttg tctaagactt acaatgttga tagacaagtt 60
263 ccagattctg c
265 <210> SEQ ID NO: 11
266 <211> LENGTH: 71
267 <212> TYPE: DNA
268 <213> ORGANISM: Artificial Sequence
270 <220> FEATURE:
271 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificial
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275 tctggaactt g
277 <210> SEQ ID NO: 12
278 <211> LENGTH: 72
279 <212> TYPE: DNA
280 <213> ORGANISM: Artificial Sequence
282 <220> FEATURE:
283 <223> OTHER INFORMATION: Description of Artificial Sequence: Artificial
285 <400> SEQUENCE: 12
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287 gaggtaatga ag
289 <210> SEQ ID NO: 13
290 <211> LENGTH: 74
291 <212> TYPE: DNA
292 <213> ORGANISM: Artificial Sequence
294 <220> FEATURE:
295 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
297 <400> SEQUENCE: 13
298 agtaacaaca ccaacagcct taccagcctt cttagctcta ttaataacag aagtaacttc 60
299 attacctcta gtag
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301 <210> SEQ ID NO: 14
302 <211> LENGTH: 74
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307 <223> OTHER INFORMATION: Description of Artificial Sequence: (Artificial
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311 catactgtta atag
313 <210> SEQ ID NO: 15
314 <211> LENGTH: 68
315 <212> TYPE: DNA
316 <213> ORGANISM: Artificial Sequence
318 <220> FEATURE:
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VERIFICATION SUMMARY

DATE: 02/25/2002

PATENT APPLICATION: US/09/911,132

TIME: 10:07:44

Input Set : A:\seq listing.txt

Output Set: N:\CRF3\02252002\I911132.raw

L:9 M:270 C: Current Application Number differs, Replaced Application Number L:10 M:271 C: Current Filing Date differs, Replaced Current Filing Date